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IN THE CLAIMS:

- 1. An isolated polynucleotide which encodes a protein comprising the amino acid sequence of SEQ ID NO:2.
- 2. The isolated polynucleotide of Claim 1, wherein said protein has serine/threonine kinase activity.
 - 3. An isolated polynucleotide, which comprises SEQ ID NO:1.
 - 4. An isolated polynucleotide which is complimentary to the polynucleotide of Claim 3.
 - 5. An isolated polynucleotide which is at least 70% identical to the polynucleotide of Claim 3.
 - 6. An isolated polynucleotide which is at least 80% identical to the polynucleotide of Claim 3.
 - 7. An isolated polynucleotide which is at least 90% identical to the polynucleotide of Claim 3.
 - 8. An isolated polynucleotide which hybridizes under stringent conditions to the polynucleotide of Claim 3; wherein said stringent conditions comprise washing in 5X SSC at a temperature from 50 to 68°C.
 - 9. The isolated polynucleotide of Claim 3, which encodes a protein having serine/threonine kinase activity.
 - 10. A vector comprising the isolated polynucleotide of Claim 1.
 - 11. A vector comprising the isolated polynucleotide of Claim 3.
 - 12. A host cell comprising the isolated polynucleotide of Claim 1.
 - 13. A host cell comprising the isolated polynucleotide of Claim 3.
 - 14. A plant cell comprising the isolated polynucleotide of Claim 1.
 - 15. A plant cell comprising the isolated polynucleotide of Claim 3.

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- 16. A transgenic plant comprising the isolated polynucleotide sequence of Claim 1.
- 17. A transgenic plant comprising the isolated polynucleotide sequence of Claim 3.
- 18. The transgenic plant of Claim 16, wherein said plant is Arabidopsis thaliania.
- 19. The transgenic plant of Claim 17, wherein said plaint is Arabidopsis thaliania.
- 20. The transgenic plant of Claim 16, wherein said plant is selected from the group consisting of wheat, corn, peanut cotton, oat, and soybean plant.
- 21. The transgenic plant of Claim 16, wherein the isolated polynucleotide is operably linked to an inducible promoter.
- 22. The transgenic plant of Claim 17, wherein the isolated polynucleotide is operably linked to an inducible promoter.
- 23. A process for screening for polynucleotides which encode a protein having serine/threonine kinase activity comprising hybridizing the isolated polynucleotide of Claim 1 to the polynucleotide to be screened; expressing the polynucleotide to produce a protein; and detecting the presence or absence of serine/threonine kinase activity in said protein.
- 24. A process for screening for polynucleotides which encode a protein having serine/threonine kinase activity comprising hybridizing the isolated polynucleotide of Claim 3 to the polynucleotide to be screened; expressing the polynucleotide to produce a protein; and detecting the presence or absence of serine/threonine kinase activity in said protein.
- 25. A process for screening for polynucleotides which encode a protein having serine/threonine kinase activity comprising hybridizing the isolated polynucleotide of Claim 8 to the polynucleotide to be screened; expressing the polynucleotide to produce a protein; and detecting the presence or absence of serine/threonine kinase activity in said protein.
- 26. A method for detecting a nucleic acid with at least 70% homology to nucleotide of Claim 1, comprising contacting a nucleic acid sample with a probe or primer comprising at least 15 consecutive nucleotides of the nucleotide sequence of Claim 1, or at least 15 consecutive nucleotides of the complement thereof.
 - 27. A method for producing a nucleic acid with at least 70% homology to nucleotide of

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Claim 1, comprising contacting a nucleic acid sample with a primer comprising at least 15 consecutive nucleotides of the nucleotide sequence of Claim 1, or at least 15 consecutive nucleotides of the complement thereof.

- 28. A method for detecting a nucleic acid with at least 70% homology to nucleotide of Claim 3, comprising contacting a nucleic acid sample with a probe or primer comprising at least 15 consecutive nucleotides of the nucleotide sequence of Claim 3, or at least 15 consecutive nucleotides of the complement thereof.
- 29. A method for producing a nucleic acid with at least 70% homology to nucleotide of Claim 3, comprising contacting a nucleic acid sample with a primer comprising at least 15 consecutive nucleotides of the nucleotide sequence of Claim 3, or at least 15 consecutive nucleotides of the complement thereof.
- 30. A method for making SOS2 protein, comprising culturing the host cell of Claim 12 for a time and under conditions suitable for expression of SOS2, and collecting the SOS2 protein.
- 31. A method for making SOS2, comprising culturing the host cell of Claim 13 for a time and under conditions suitable for expression of SOS2, and collecting the SOS2 protein.
- 32. A method of making a transgenic plant comprising introducing the polynucleotide of Claim 1 into the plant.
- 33. A method of making a transgenic plant comprising introducing the polynucleotide of Claim 1 into the plant.
- 34. A method of increasing the salt tolerance of a plant in need thereof, comprising introducing the polynucleotide of Claim 1 into said plant.
- 35. A method of increasing the salt tolerance of a plant in need thereof, comprising introducing the polynucleotide of Claim 1 into said plant.
- 36. A method of increasing the salt tolerance of a plant in need thereof, comprising enhancing the expression of the SOS 2 gene into said plant.
 - 37. An isolated polypeptide comprising the amino acid sequence in SEQ ID NO:2.

- 38. The isolated polypeptide of Claim 37 which has serine/threonine kinase activity.
- 39. An isolated polypeptide which is at least 70% identical to the isolated polypeptide of Claim 37 and which has serine/threonine kinase activity.
- 40. An isolated polypeptide which is at least 80% identical to the isolated polypeptide of Claim 37 and which has serine/threonine kinase activity.
- 41. An isolated polypeptide which is at least 90% identical to the isolated polypeptide of Claim 37 and which has serine/threonine kinase activity.
- 42. An isolated polypeptide which is at least 95% identical to the isolated polypeptide of Claim 37 and which has serine/threonine kinase activity.